## **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

10/5/8,938
PCT/10
1/10/05

## ENTERED



PCT

DATE: 01/10/2005 RAW SEQUENCE LISTING TIME: 15:42:25 PATENT APPLICATION: US/10/518,938

Input Set : A:\82368-18.app

```
3 <110> APPLICANT: Nakamura, Yusuke
             Furukawa, Yoichi
             Oncotherapy Science, Inc.
             The University of Tokyo
      8 <120> TITLE OF INVENTION: Method for Diagnosis of Colorectal Tumors
     10 <130> FILE REFERENCE: 082368-001800US
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/518,938
C--> 13 <141> CURRENT FILING DATE: 2004-12-17
    15 <150> PRIOR APPLICATION NUMBER: US 60/389,994
   16 <151> PRIOR FILING DATE: 2002-06-19
     18 <150> PRIOR APPLICATION NUMBER: WO PCT/JP02/12760
     19 <151> PRIOR FILING DATE: 2002-12-05
     21 <160> NUMBER OF SEQ ID NOS: 21
    23 <170> SOFTWARE: PatentIn version 3.1
     25 <210> SEQ ID NO: 1
     26 <211> LENGTH: 27
     27 <212> TYPE: DNA
     28 <213> ORGANISM: Artificial
     30 <220> FEATURE:
     31 <223> OTHER INFORMATION: artificially synthesized primer sequence
     33 <400> SEQUENCE: 1
                                                                               27
     34 tccctgttgt aatcaagaat aagacct
     37 <210> SEQ ID NO: 2
     38 <211> LENGTH: 25
     39 <212> TYPE: DNA
     40 <213> ORGANISM: Artificial
     42 <220> FEATURE:
     43 <223> OTHER INFORMATION: artificially synthesized primer sequence
     45 <400> SEQUENCE: 2
                                                                               25
     46 caqttccttt gcagtaagat ccatc
     49 <210> SEQ ID NO: 3
     50 <211> LENGTH: 36
     51 <212> TYPE: DNA
     52 <213> ORGANISM: Artificial
     54 <220> FEATURE:
     55 <223> OTHER INFORMATION: artificially synthesized probe sequence
     57 <220> FEATURE:
     58 <221> NAME/KEY: misc_binding
     59 <222> LOCATION: (1)..(1)
     60 <223> OTHER INFORMATION: labeled with VIC
     62 <220> FEATURE:
     63 <221> NAME/KEY: misc_binding
     64 <222> LOCATION: (36)..(36)
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Input Set : A:\82368-18.app

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65 <223> OTHER INFORMATION: labeled with TAMRA
67 <400> SEQUENCE: 3
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68 ttgttgaagg tctccctatt aatgatttct cacgtg
71 <210> SEQ ID NO: 4
72 <211> LENGTH: 21
73 <212> TYPE: DNA
74 <213> ORGANISM: Artificial
76 <220> FEATURE:
77 <223> OTHER INFORMATION: artificially synthesized primer sequence
79 <400> SEQUENCE: 4
                                                                           21
80 gcagactctg cgcttgagat c
83 <210> SEQ ID NO: 5
84 <211> LENGTH: 20
85 <212> TYPE: DNA
86 <213> ORGANISM: Artificial
88 <220> FEATURE:
89 <223> OTHER INFORMATION: artificially synthesized primer sequence
91 <400> SEQUENCE: 5
                                                                           20
92 gggctagtcg cacagacctc
95 <210> SEQ ID NO: 6
96 <211> LENGTH: 26
97 <212> TYPE: DNA
98 <213> ORGANISM: Artificial
100 <220> FEATURE:
101 <223> OTHER INFORMATION: artificially synthesized probe sequence
103 <220> FEATURE:
104 <221> NAME/KEY: misc binding
105 <222> LOCATION: (1)..(1)
106 <223> OTHER INFORMATION: labeled with FAM
108 <220> FEATURE:
109 <221> NAME/KEY: misc binding
110 <222> LOCATION: (26)..(26)
111 <223> OTHER INFORMATION: labeled with TAMRA
113 <400> SEQUENCE: 6
                                                                            26
114 aacaagcatc agcgttttcc agggct
117 <210> SEQ ID NO: 7
118 <211> LENGTH: 19
119 <212> TYPE: DNA
120 <213> ORGANISM: Artificial
122 <220> FEATURE:
123 <223> OTHER INFORMATION: artificially synthesized primer sequence
125 <400> SEQUENCE: 7
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126 caaatggctg ccaaactgg
129 <210> SEQ ID NO: 8
130 <211> LENGTH: 22
131 <212> TYPE: DNA
132 <213> ORGANISM: Artificial
134 <220> FEATURE:
135 <223> OTHER INFORMATION: artificially synthesized primer sequence
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Input Set : A:\82368-18.app

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137 <400> SEQUENCE: 8
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138 ggattctttg ttcttccgca ct
141 <210> SEQ ID NO: 9
142 <211> LENGTH: 31
143 <212> TYPE: DNA
144 <213> ORGANISM: Artificial
146 <220> FEATURE:
147 <223> OTHER INFORMATION: artificially synthesized probe sequence
149 <220> FEATURE:
150 <221> NAME/KEY: misc_binding
151 <222> LOCATION: (1)..(1)
152 <223> OTHER INFORMATION: labeled with FAM
154 <220> FEATURE:
155 <221> NAME/KEY: misc_binding
156 <222> LOCATION: (31)..(31)
157 <223> OTHER INFORMATION: labeled with TAMRA
159 <400> SEQUENCE: 9
                                                                            31
160 cqtttqcgag agaaggataa gcacattgaa g
163 <210> SEQ ID NO: 10
164 <211> LENGTH: 24
165 <212> TYPE: DNA
166 <213> ORGANISM: Artificial
168 <220> FEATURE:
169 <223> OTHER INFORMATION: artificially synthesized primer sequence
171 <400> SEQUENCE: 10
                                                                            24
172 gacagcagtg gagaattgat gttt
175 <210> SEQ ID NO: 11
176 <211> LENGTH: 26
177 <212> TYPE: DNA
178 <213> ORGANISM: Artificial
180 <220> FEATURE:
181 <223> OTHER INFORMATION: artificially synthesized primer sequence
183 <400> SEQUENCE: 11
                                                                            26
184 acaatttgag gacacttcat atttgc
187 <210> SEQ ID NO: 12
188 <211> LENGTH: 22
189 <212> TYPE: DNA
190 <213> ORGANISM: Artificial
192 <220> FEATURE:
193 <223> OTHER INFORMATION: artificially synthesized probe sequence
195 <220> FEATURE:
196 <221> NAME/KEY: misc binding
197 <222> LOCATION: (1)..(1)
198 <223> OTHER INFORMATION: labeled with FAM
200 <220> FEATURE:
201 <221> NAME/KEY: misc_binding
202 <222> LOCATION: (22)..(22)
203 <223> OTHER INFORMATION: labeled with TAMRA
205 <400> SEQUENCE: 12
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Input Set : A:\82368-18.app

		22
	tgaggcagac ttggtgctgg cg	44
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	<211> LENGTH: 24	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial	
214	<220> FEATURE:	
	<223> OTHER INFORMATION: artificially synthesized primer sequence	
	<400> SEQUENCE: 13	0.4
	gcatacaagt tggcaggaac atta	24
	<210> SEQ ID NO: 14	
	<211> LENGTH: 21	
	<212> TYPE: DNA	
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	<220> FEATURE:	
227	<223> OTHER INFORMATION: artificially synthesized primer sequence	
229	<400> SEQUENCE: 14	
230	accacaagec gateteette t	21
233	<210> SEQ ID NO: 15	
234	<211> LENGTH: 27	
	<212> TYPE: DNA	
236	<213> ORGANISM: Artificial	
	<220> FEATURE:	
239	<223> OTHER INFORMATION: artificially synthesized probe sequence	
241	<220> FEATURE:	
	<221> NAME/KEY: misc_binding	
	<222> LOCATION: (1)(1)	
244	<223> OTHER INFORMATION: labeled with FAM	
246	<220> FEATURE:	
	<221> NAME/KEY: misc_binding	
248	<222> LOCATION: (27)(27)	
249	<223> OTHER INFORMATION: labeled with TAMRA	
251	<400> SEQUENCE: 15	
252	catggcagtg attctgtgga gagtgca	27
255	<210> SEQ ID NO: 16	
256	<211> LENGTH: 21	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial	
260	<220> FEATURE:	
	<223> OTHER INFORMATION: artificially synthesized primer sequence	
263	<400> SEQUENCE: 16	
	agatgcggaa aaacttgacc a	21
267	<210> SEQ ID NO: 17	
	<211> LENGTH: 20	
	<212> TYPE: DNA	
	<213> ORGANISM: Artificial	
272	<220> FEATURE:	
273	<223> OTHER INFORMATION: artificially synthesized primer sequence	
	<400> SEQUENCE: 17	
276	agtcagtctg ggtcccacca	20

Input Set : A:\82368-18.app

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279 <210> SEQ ID NO: 18
280 <211> LENGTH: 26
281 <212> TYPE: DNA
282 <213> ORGANISM: Artificial
284 <220> FEATURE:
285 <223> OTHER INFORMATION: artificially synthesized probe sequence
287 <220> FEATURE:
288 <221> NAME/KEY: misc_binding
289 <222> LOCATION: (1)..(1)
290 <223> OTHER INFORMATION: labeled with FAM
292 <220> FEATURE:
293 <221> NAME/KEY: misc_binding
294 <222> LOCATION: (26)..(26)
295 <223> OTHER INFORMATION: labeled with TAMRA
297 <400> SEQUENCE: 18
                                                                            26
298 aagccatcag agagcatcag atggcc
301 <210> SEQ ID NO: 19
302 <211> LENGTH: 20
303 <212> TYPE: DNA
304 <213> ORGANISM: Artificial
306 <220> FEATURE:
307 <223> OTHER INFORMATION: artificially synthesized primer sequence
309 <400> SEQUENCE: 19
                                                                            20
310 gcagcgttat acgcagagca
313 <210> SEQ ID NO: 20
314 <211> LENGTH: 19
315 <212> TYPE: DNA
316 <213> ORGANISM: Artificial
318 <220> FEATURE:
319 <223> OTHER INFORMATION: artificially synthesized primer sequence
321 <400> SEQUENCE: 20
                                                                            19
322 acccacgatg agggcagag
325 <210> SEQ ID NO: 21
326 <211> LENGTH: 18
327 <212> TYPE: DNA
328 <213> ORGANISM: Artificial
330 <220> FEATURE:
331 <223> OTHER INFORMATION: artificially synthesized probe sequence
333 <220> FEATURE:
 334 <221> NAME/KEY: misc_binding
 335 <222> LOCATION: (1)..(1)
 336 <223> OTHER INFORMATION: labeled with FAM
 338 <220> FEATURE:
 339 <221> NAME/KEY: misc_binding
 340 <222> LOCATION: (18)..(18)
 341 <223> OTHER INFORMATION: labeled with TAMRA
 343 <400> SEQUENCE: 21
                                                                             18
 344 tgggcgcagg ccgtttgg
```

Input Set : A:\82368-18.app

Output Set: N:\CRF4\01102005\J518938.raw

## Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21

VERIFICATION SUMMARY

DATE: 01/10/2005

PATENT APPLICATION: US/10/518,938

TIME: 15:42:26

Input Set : A:\82368-18.app

Output Set: N:\CRF4\01102005\J518938.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application Number L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date